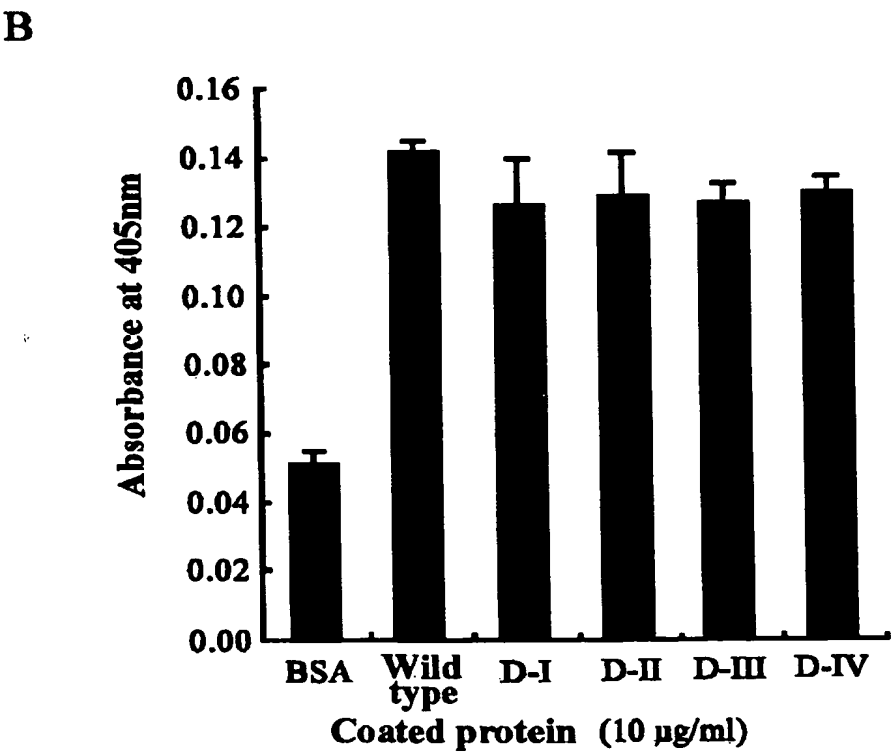
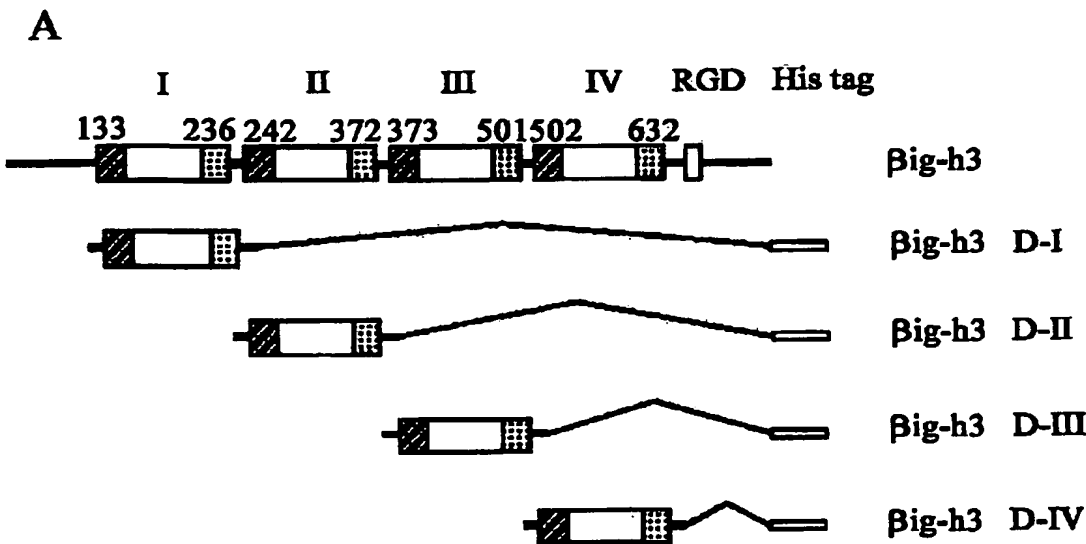


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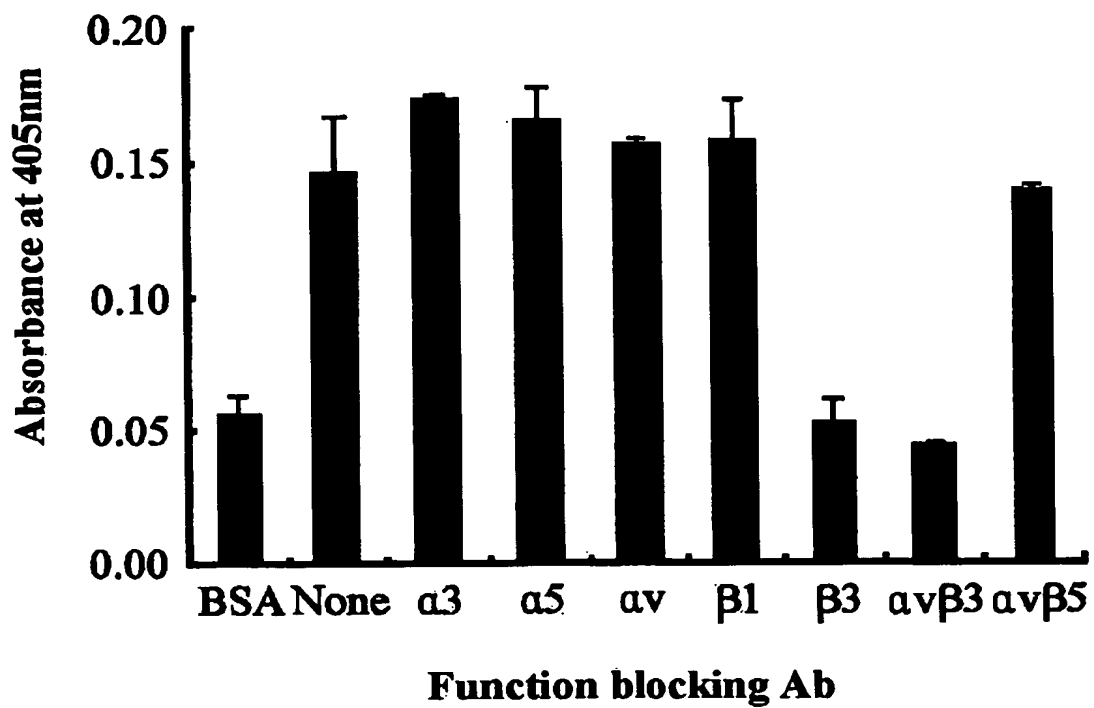
FIG. 1

BIGH3_HUMAN	161	SNVNIELINAEIRYH	MVGRRV	PTDEPKHGMT	190					
BIGH3-PIG	161	SNVNIELINAEIRYH	MVDRRV	PTDEPKHGMA	190					
BIGH3_CHICK	154	SNVNIELINAEIRYH	MVNKRV	PTDEPKHGTT	183					
OSF2_MOUSE	157	NNVNVELINAEIRYH	MVNKRV	PTDEPKHGMV	186					
BIGH3_HUMAN	298	-GDPEARLDLENNH	IKSAMCAE	IVAGLS	326					
BIGH3-PIG	298	-GDPEARLDLENNH	IKSAMCAE	IVAGLS	326					
BIGH3_CHICK	290	-GDPEARLDLENNH	IKSAMCAE	IVAGLT	318					
OSF2_HUMAN	292	-GDKVASEAEMKYH	INTLQCSE	IVMGGAV	320					
OSF2_MOUSE	294	-GDKVASEAEMKYH	INTLQCSE	IVTGGAV	322					
BIGH3_HUMAN	560	-GDAKELANEIRKYH	IGDEIVSGG	IGALVR	588					
BIGH3-PIG	560	-GNAKELANEIRKYH	VGDEIVSGG	IGALVR	588					
BIGH3_CHICK	552	-GNAKELANEIRKFH	MADENIVSG	AVSALVR	580					
SLL1735 homolog	59	-QNPPQAFELITYH	VAAGH	ITKDD	IFIGE 87					
SLL1735	59	-QNIPQAFELITYH	VVAGKFTQAD	CHIST	87					
SLL1483	104	PENKDFLVFELITYH	VVPGR	ITAAQV-QSGE	132					
OSF2_HUMAN	554	-RDKNALQNEELITYH	ETPGVF	IGKGFEPGVT	582					
OSF2_MOUSE	556	-GDKNALQNEELITYH	ETPGVF	IGKGFEPGVT	584					
MP83 MYCTU	164	-TDAKLESEELITYH	VEAGQ--	ASPSRIDGT	190					
MPT83	145	-TDAKLESEELITYH	VEAGQ--	ASPSRIDGT	171					
Q48948_MYCBO	119	-TNSSLETSLELITYH	VVAGQ--	TSPANVVGT	145					
Q50769_MYCTU	119	-TNSSLETSLELITYH	VVAGQ--	TSPANVVGT	145					
Putative Secreted protein	132	-NDRAQPKKVELITYH	VVEHKR	ITKAC	IPHGT 160					
Fasciclin	485	EG-RGCASNEIRK	NH	LDLTFC	SLATVPGAK 513					
HLC-32	284	KDPAGHERNLEIRKYH	V	SDVKYSV	LSGQR 313					
Fasciclin	775	-SKPADPMELVK	TH	VEDDVCCAG	IPPTNW 803					
BIGH3_HUMAN	432	RNLIR.NH	IKDQ	ASKY	LYH	GQT	DT	IGGK	IR	464
BIGH3-PIG	432	RNLIR.NH	IKDQ	ASKY	LYH	GQT	DT	IGGK	IR	464
BIGH3_CHICK	424	RNLIR.NH	IKDQ	SSKY	LYH	GQT	DT	IGGK	IR	456
OSF2_HUMAN	426	-RNLIR.NH	IKDQ	SSKY	LYH	GQT	DT	IGGK	IR	458
OSF2_MOUSE	428	-RNLIR.NH	IKDQ	SSKY	LYH	GQT	DT	IGGK	IR	460

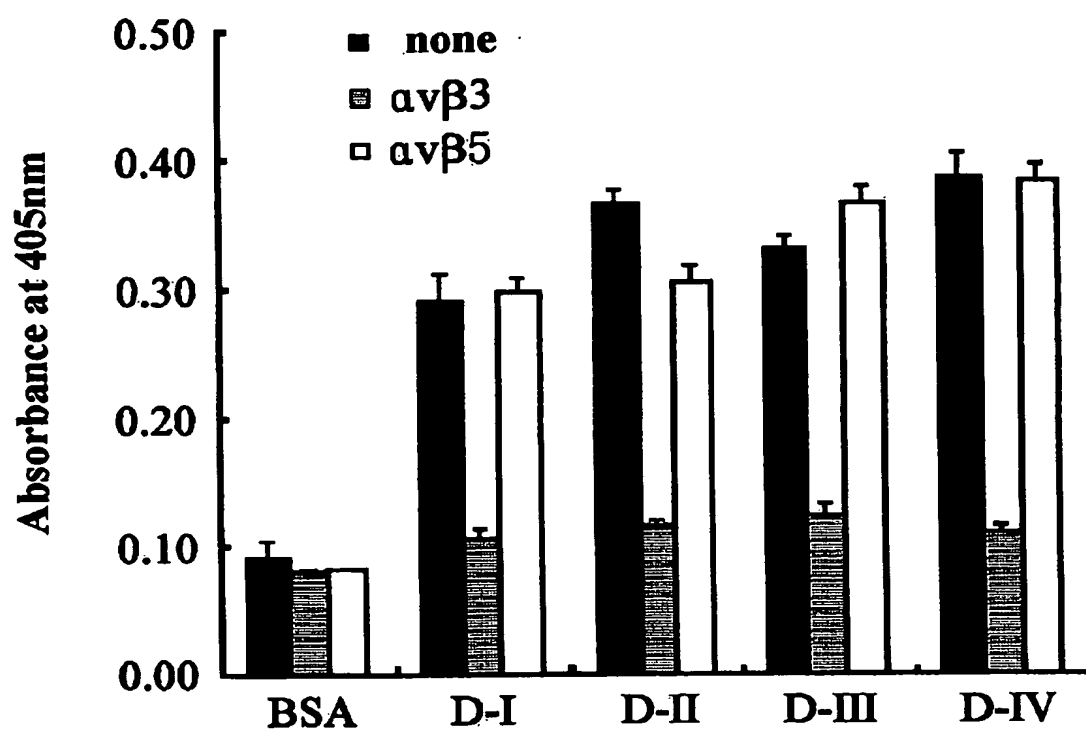
FIG. 2



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FIG. 3a

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FIG. 3b

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FIG. 3c

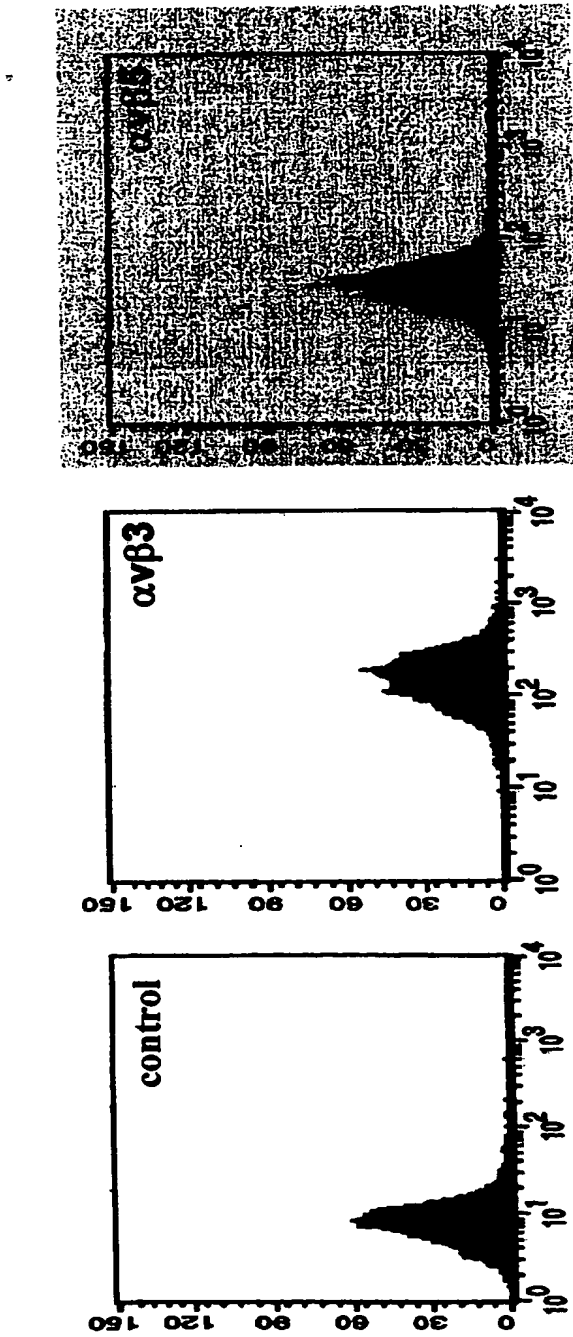
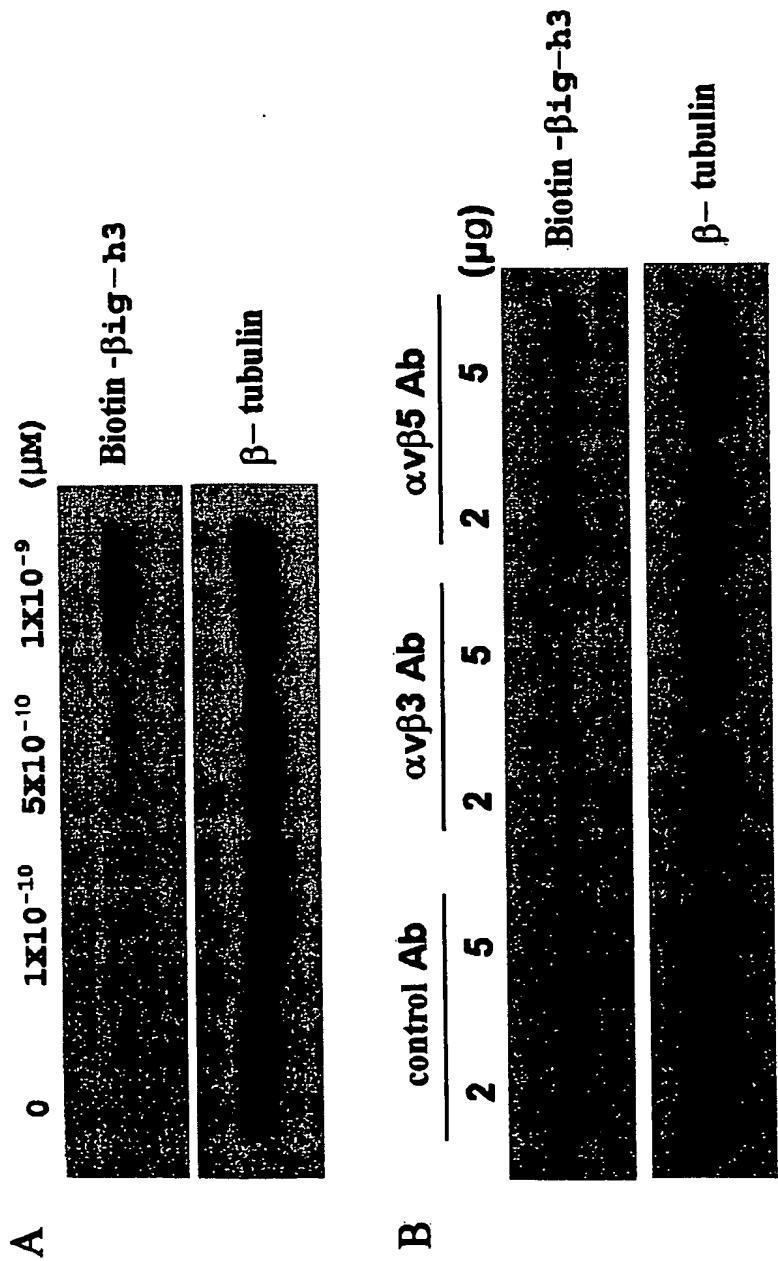


FIG. 3d



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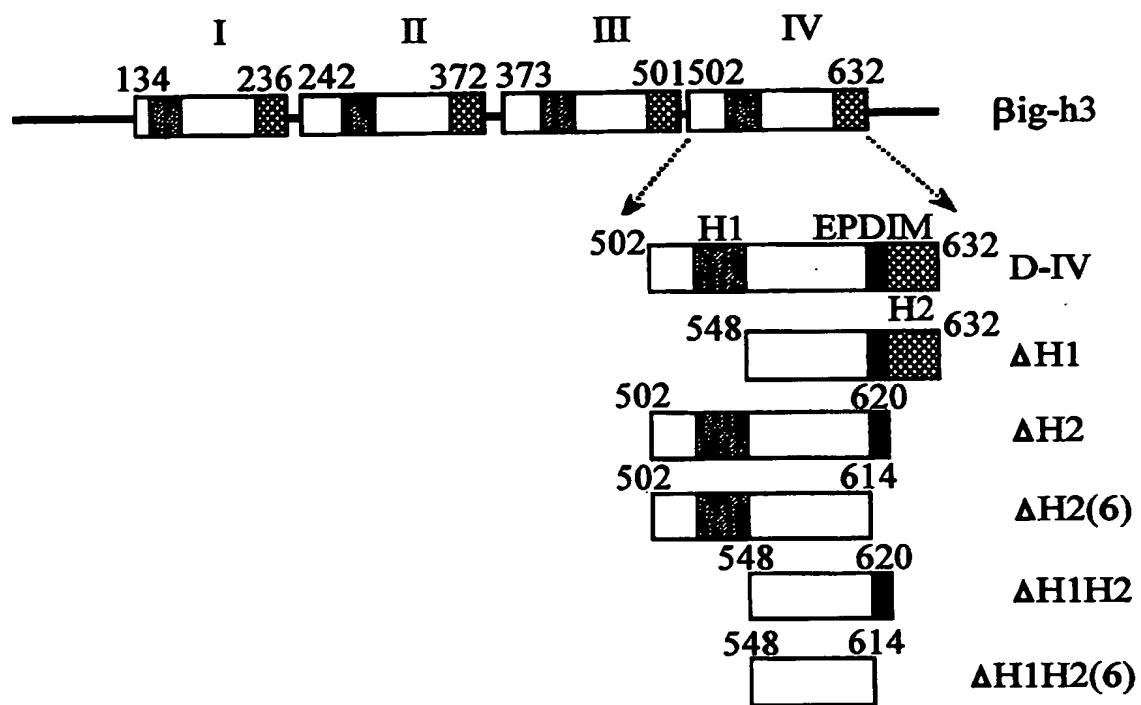
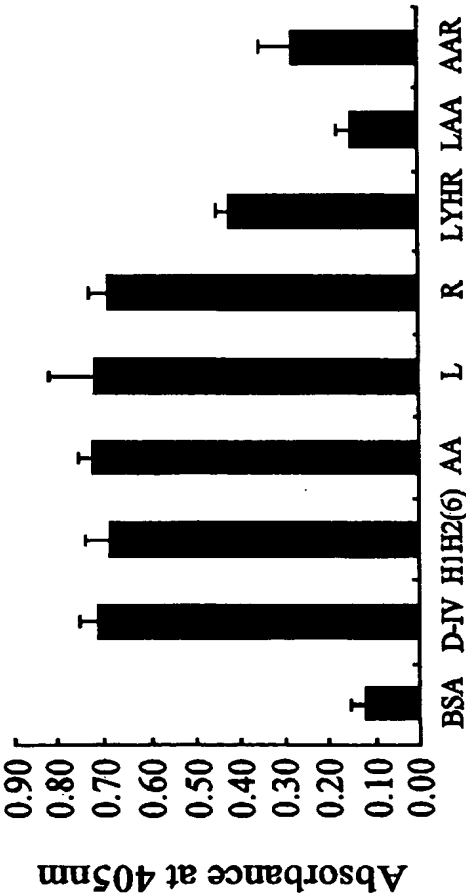
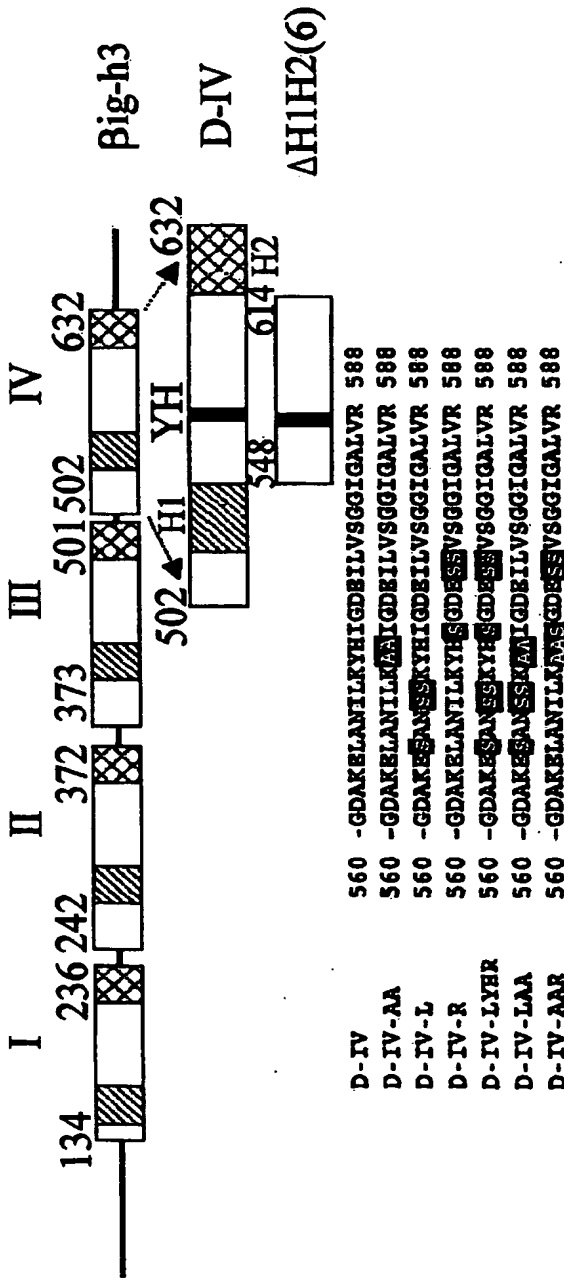
FIG. 4a

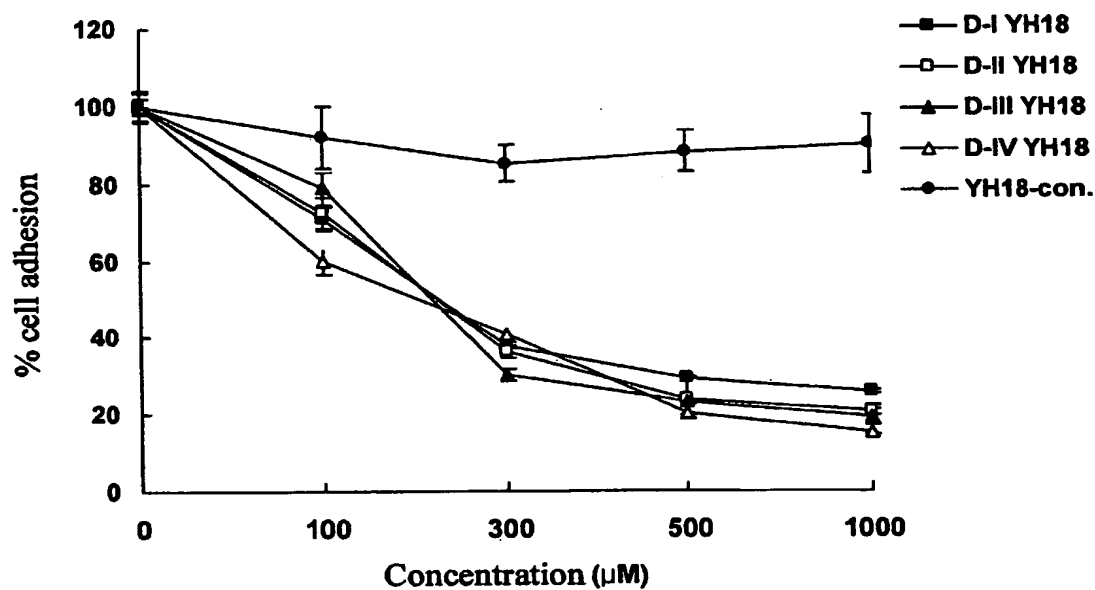
FIG. 4b



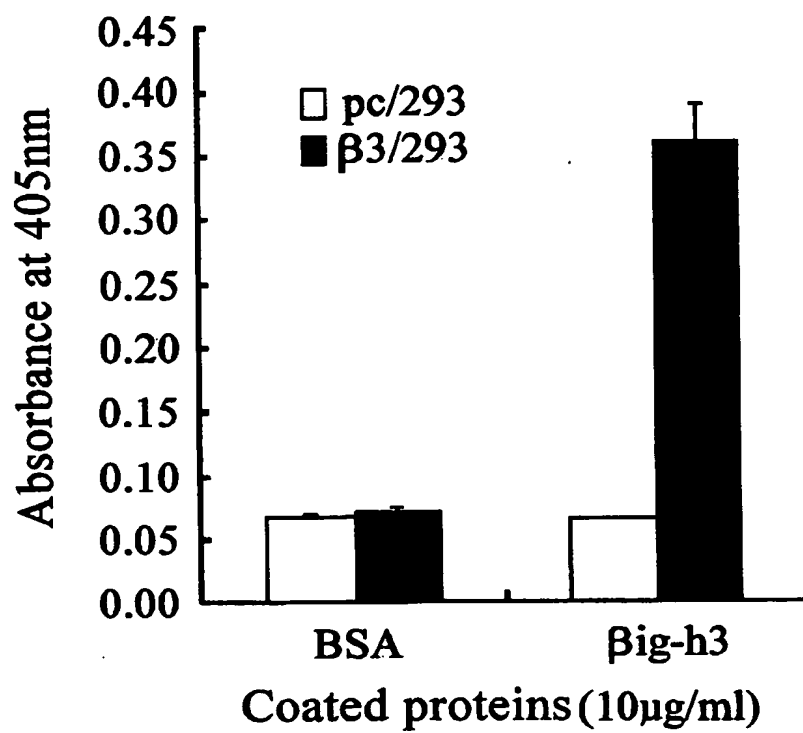
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FIG. 5

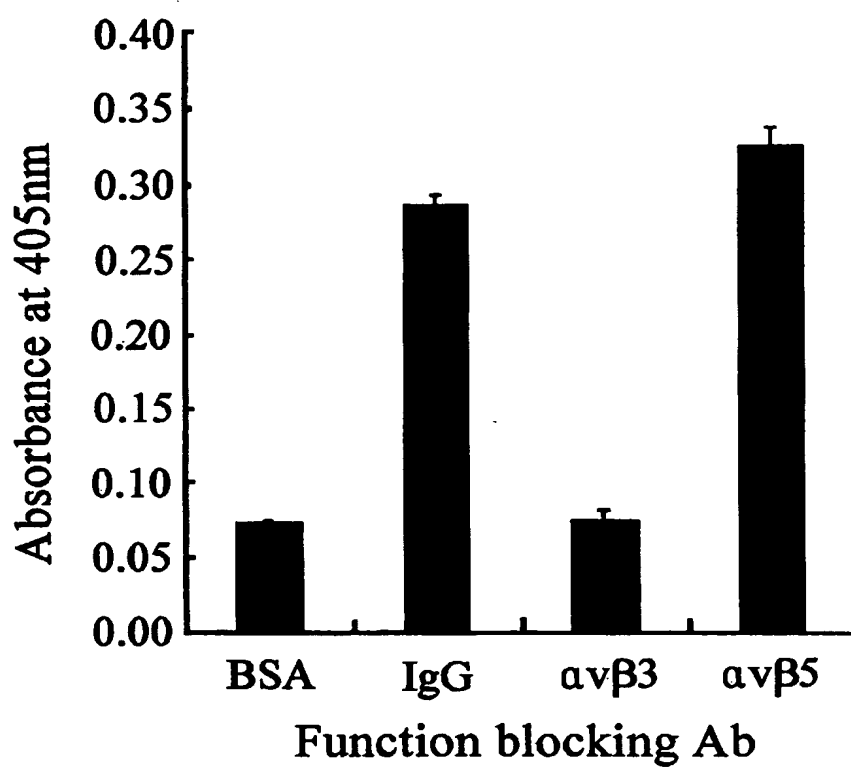
D-I	YH18	165	IELLNALRYEMVGRRVLT	182
D-II	YH18	301	EALRDLNNHILKSAMCA	318
D-III	YH18	442	DQLASKYLYHGQTLETG	459
D-IV	YH18	563	KELANILKYHIGDEILVS	580
YH18-con			KELANIHGKLYDEILVS	



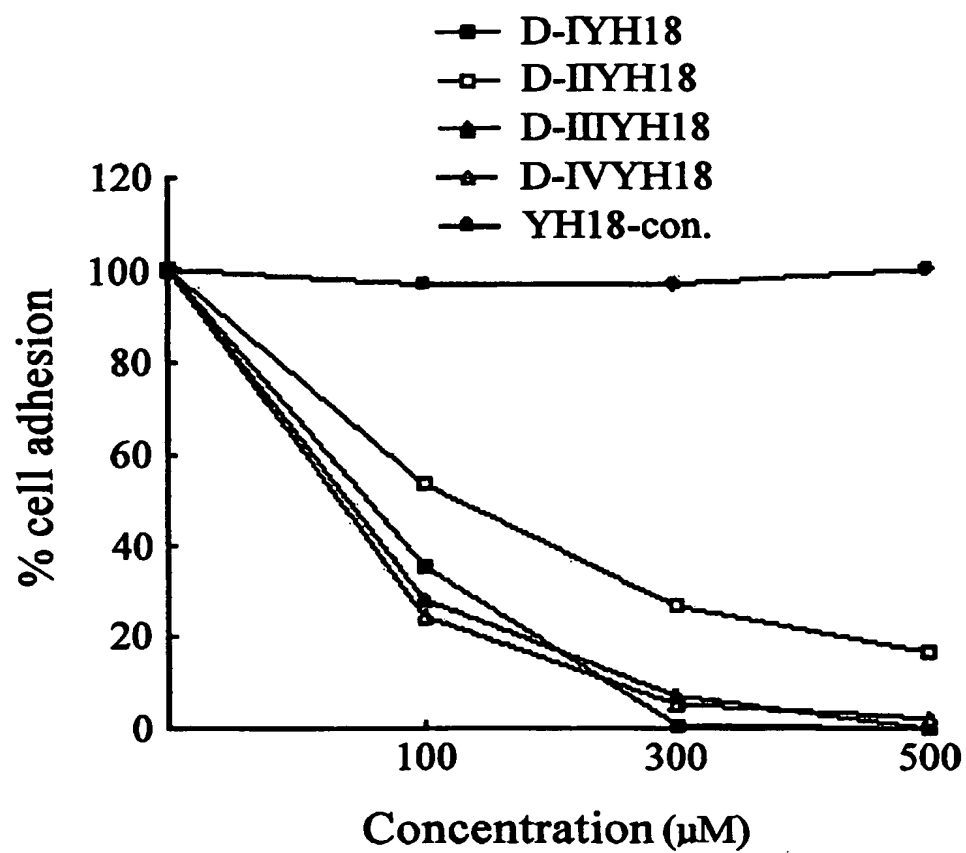
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FIG. 6a

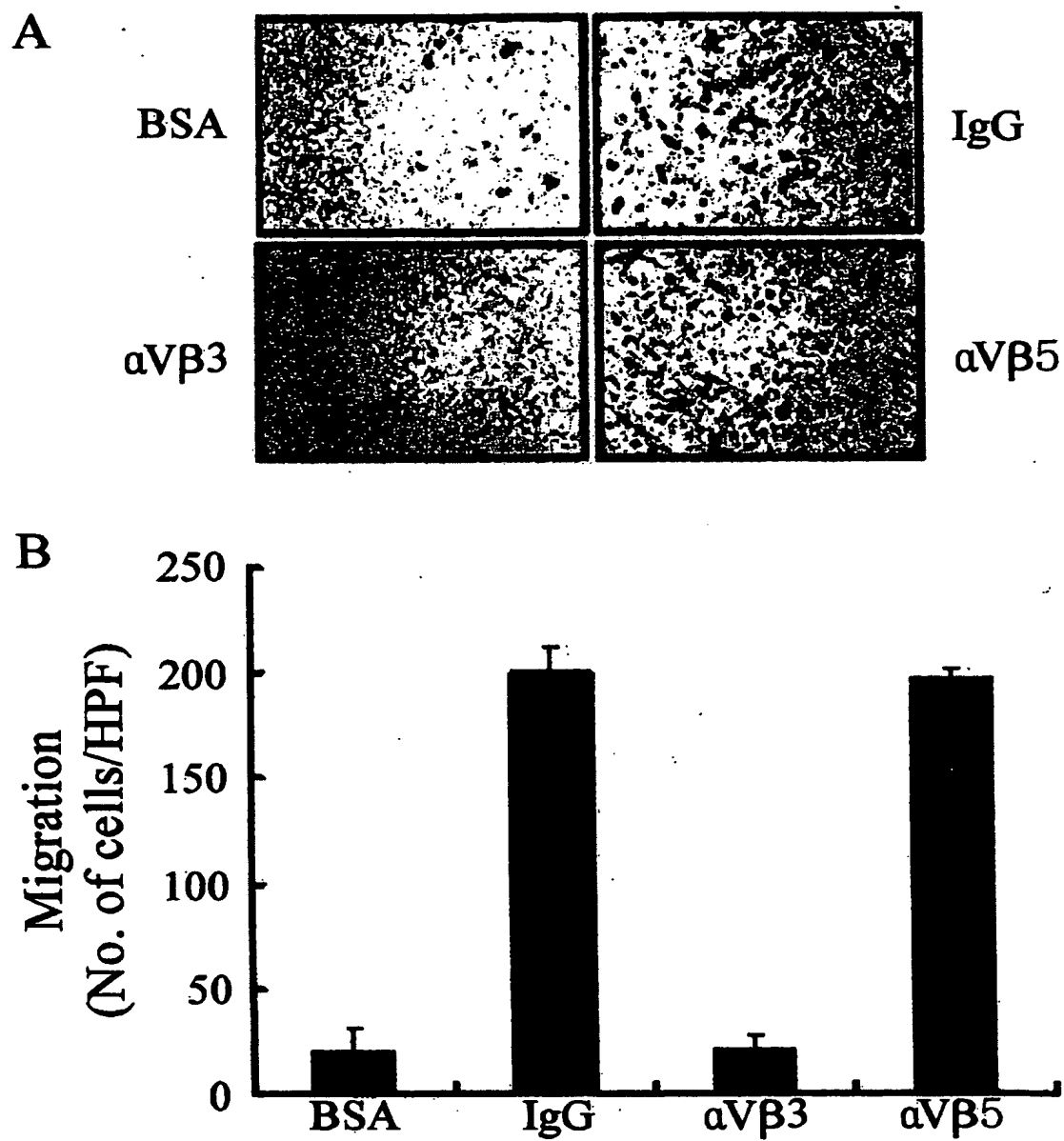
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FIG. 6b

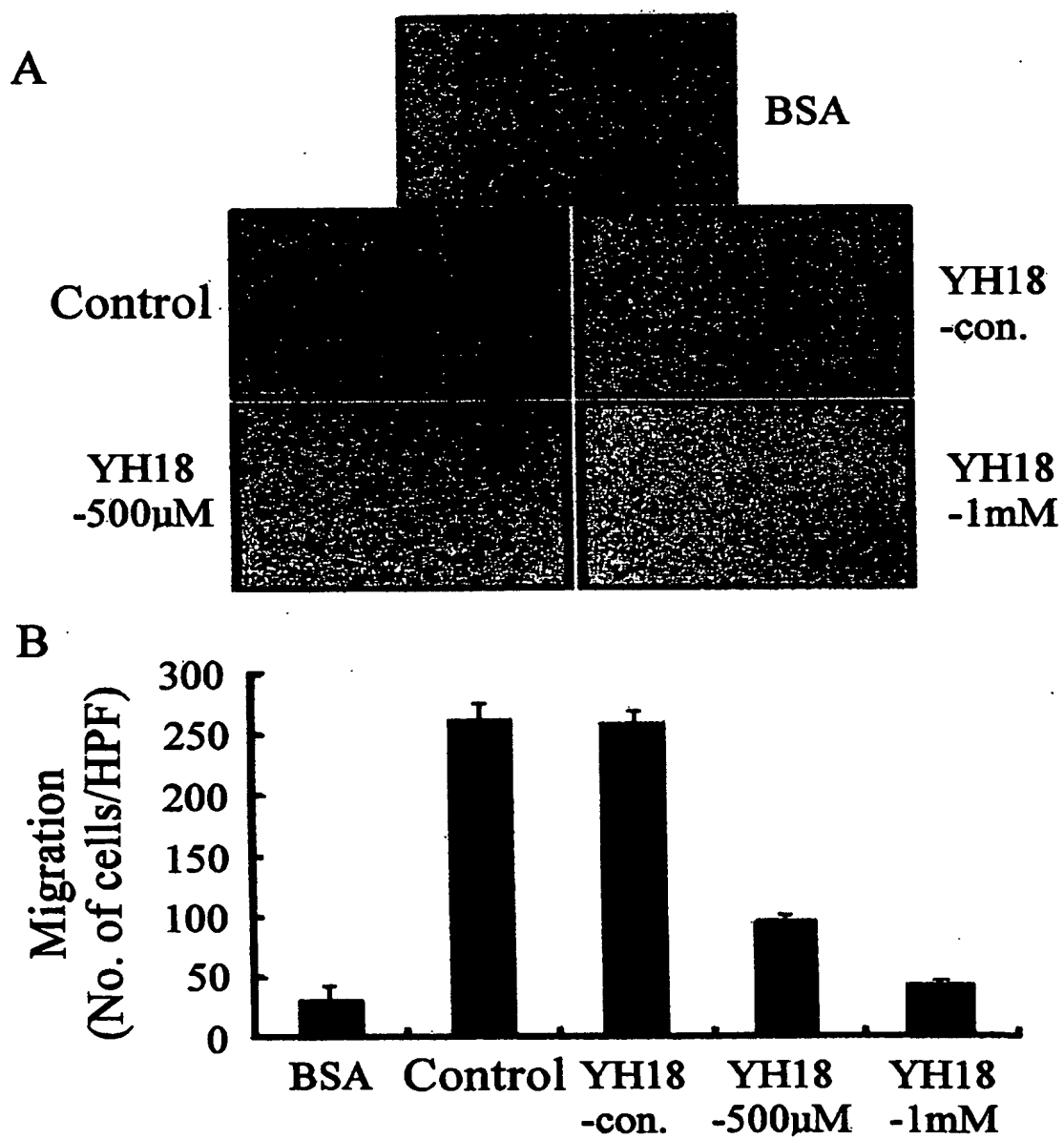
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FIG. 6c

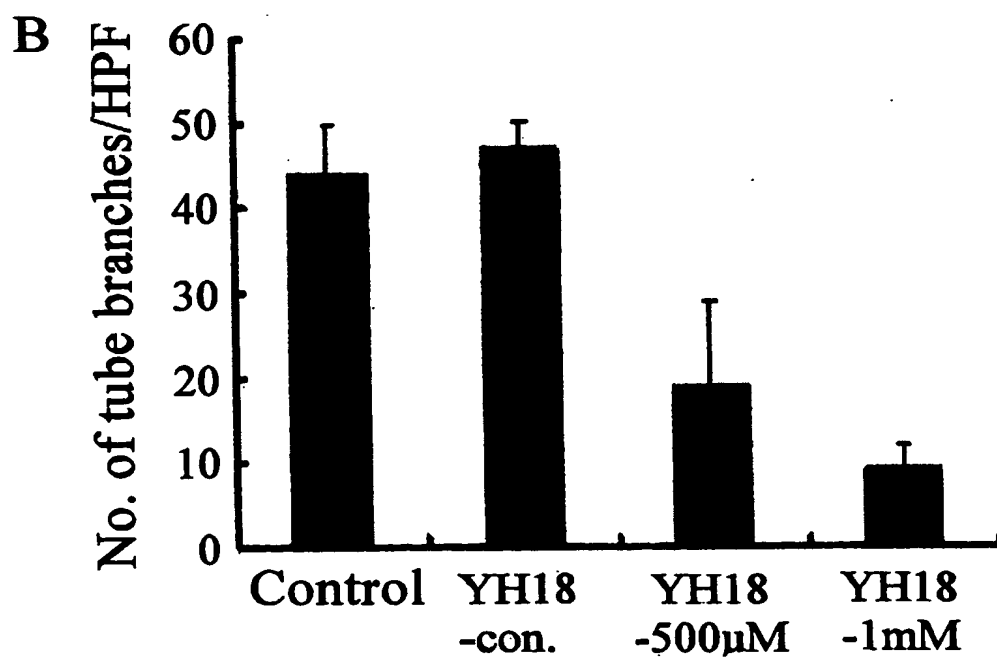
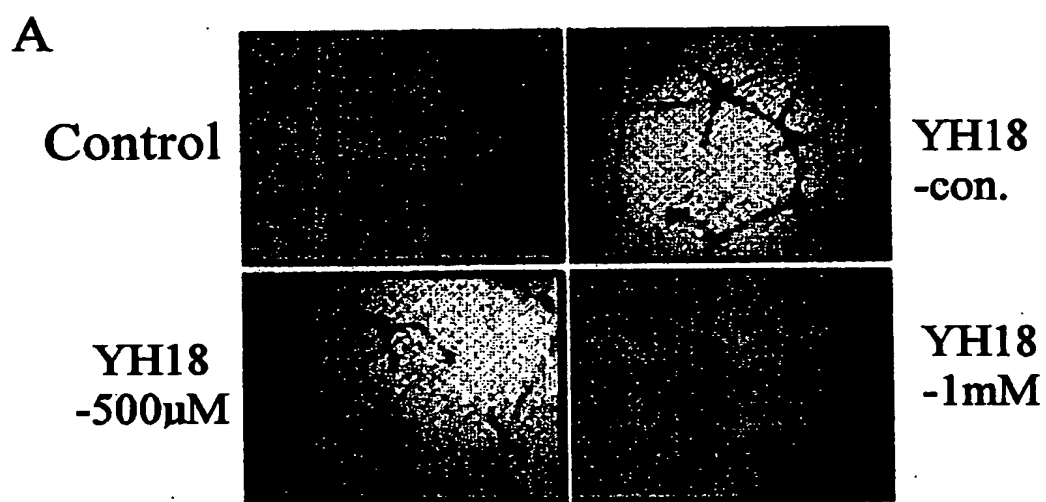
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FIG. 7a

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FIG. 7b

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FIG. 8a

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FIG. 8b